SUBJECT: Abbreviated Construct Hazard Assessment for J15-31 FROM: Gwendolyn McClung, Ph.D. Technical Integrator Assessment Branch 3 Risk Assessment Division (7403M) THRU: Cathy Fehrenbacher, CIH Chief Assessment Branch 3 Risk Assessment Division (7403M) DATE: July 30, 2015 I. INTRODUCTION The Agency has received a Microbial Commercial Activity Notice (MCAN) from Danisco US, Inc. (operating as DuPont Industrial Biosciences) for an intergeneric Saccharomyces cerevisiae strain that has In a previous submission the recipient *S. cerevisiae* strain was modified by the introduction of a In the current submission, a resulting in the new submission strain . The parental strain used in fuel ethanol strain is S. cerevisiae production that has that can then be fermented into ethanol by the microorganism. The production microorganism, strain , will thus be used for production of fuel ethanol from grain. II. POTENTIAL HAZARDS POSED BY THE GENETIC MODIFICATIONS Genetic modifications of the recipient strain to arrive at the submission strain, S. cerevisiae are described in detail by Segal (2015) in the Genetic Construction Report. A. Inserted Genes gene to create the There is low hazard associated with the introduced submission strain of ). The introduced gene have a long history of safe use in the food industry.

common in microorganisms and	
which can then be fermented into ethanol.	
There is low hazard associated with the introduced gene to create submission strain <i>S. cerevisiae</i>	the
. This enz	yme is
and is also found in humans and other mammals. The	
microorganism, , is ubiquitous in the environment in soils, water, and vegetation. It has a long history of safe use for	i

## B. Potential for Horizontal Gene Transfer

The introduced genes were both stably inserted into the chromosome which means they are unlikely to horizontally transfer to other microorganisms in the environment if the submission microorganism was inadvertently released from the manufacturing facility or ethanol production plants. The scientific literature suggests that horizontal gene transfer (HGT) between fungi is low even though recently it has been suggested that HGT of genes from other organisms into fungi has been shown to be important in the evolution of fungi (Rosewich et al., 2000; Fitzpatrick, 2004; Richards et al., 2011). There is low concern for horizontal gene transfer, and there would be low concern even if the genes were to be transferred.

## III. CONCLUSIONS

There are low hazards associated with the genetic modifications done to create the
submission strains, S. cerevisiae The introduced
are commonly found in many microorganisms and merely allow for the
. These do not pose hazards. Also there is low potential for
horizontal gene transfer if the submission microorganism was inadvertently introduced
into the environment

## REFERENCES

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